

## SEQUENCE LISTING

```
<110°
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      Kieffer-Higgins, Stephen G.
      Chang, Dañíong
Sadook, Grasam F.
      Webb, Michael
      McDowell, Christopher S.
(12) Methods of and Apparatus for Separating and Detecting Nucleic Acid
0130 110163.132032
(11) THE 09/766,886
1141. . ((01-1)1-19.
<150. UN 60 (176,839)
(131. .001-)1-19
<160.8 , \oplus
(17) PastSEQ for Windows Version 4.0
-12101
H211: 10
Sulla DNA
10130 Staphylococcus epidermis
intgwar dat dtoaggroot gaoggaagda goattaagtig gatostoata
                                                                        50
-111230- 12
-11 - 16
CLIES DHA
11130 Unknown
1.230 staphylococcus
(40)00-2
anatutuagg tootgacgga agcage
                                                                         26
H2100 3
::[11: !(
:::12: FNA
HUIS Staphylococcus aureus
H14000 H
satquaccat qtcaggtcct gacggaagca gcattaagtg gatcatcata
                                                                        50
\{(1,1,1),\dots,4\}
42111-49
1212 - DNA
<215: Unknown
<223: staphylococcus
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Page i

## 11.183-18.70...

•	11.16.1-13.00	
	<pre></pre>	49
	<pre> *213</pre>	46
	Cill DNA  Cill Streptococcus pneumoniae  Cill A  yntorgogig aagogggtoa ggggaggaat beagsagoob taageg  Cill C 7  Cill C 26	46
	<pre>0.11 : DNA 0.11 : Unknown  1.2 : 0.12 : streptococcus  0.40 : 7 atgrettagg ggaggaatec ageage</pre>	26
	<pre>-001 0</pre>	46
	HULLS 46 HULLS 5NA HULLS 5NA HULLS 5NA HULLS 5NA	
	Gottigogtg aagogggtoa ggggaggaat coagoagood taagog  -(210:-10 -(.11:-41 -(.1.0:-DNA -(.11:-Unknown	46
	0.020 streptococcus  0.0200 10 ttgentgaag gggtcagggg aggaatccag cagcotaago g	41
	32112 64	

## 11 16--18 77 .....

1232 NA 1233 K.epsielia pheumoniae	
r400° 11 ogdalogota ototgtitas pagginayyt boygalaggaa goagobagag bagangalogt ytgt	. 60 64
1210 + 12 <821 + 64 1212 + DMA 813 + Unknown	
<pre>% 120 + &lt;</pre>	
<pre>&lt;400 - 12 ijoa.cqcta ototqtttac caggtcaggt onggaaggaa ghaghcaagg bagatgabg; jtqt</pre>	: €0 €4
<pre></pre>	
-(40) 13 - Montegata etetgittae eaggicaggi eeggaäggaa geageeaagg eagaigaege gig:	: 60 64
::12 = 24 :211 + 26 ::12:	
(10) - 14 4 agatoagg tooggaagga ageage	2.5
0010 + 15	
$(40) \times 10$ . Moseogatt accogtosac ctggtosggt coggasggs gosgcososg ogggsscatogigt	60 64
1010 16 00110 64 00130 DNA 00130 Bacillus cereus	
$<\!\!400$ : 16 Mgcaacggga cocgtgaacc ttgtcaggtc cggaaggaag cagcaataag cggtcttctc $_{\rm 3}$ .gc.	60 64
+.:10. 17 +.:11: 44 +.:12: DNA <213> Artificial Sequence	

## 

<220> KZZBZ diigonudledtide prope	
476 k 17 sysaasgayt anagythagy tsasyyaagga agsaysagsg gtyt	4.4
-310 - 18 <311> 20 <212 - DNA 	
<220 - 1183 - bligonuslestide probe	
1100 - 18 Lygo oggga apgtattpat	<u>2</u> 2.
CHD 19 CHL 22 CH2 DNA CH3 Artificial Sequence	
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04000-19 ottougtota otgogoacao gg	22
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